group consisting of H, cyclic aralkyl, and C_{1-6} alkyl; R_1 is a tyrosyl residue or a 2',6'-dimethyltyrosyl residue; R_2 is an amino acid having the R-configuration, aminoisobutyric acid, cyclopropylalanine, cyclohomoleucine or cycloleucine; R_3 is an aromatic amino acid; R_4 is an aromatic amino acid residue; Q is an amide bond or an interposed amide bond mimetic; with the following provisos: a) when R_1 is a tyrosyl residue; R_2 is D-alanine; X, Y, and Z, are H; and R_3 is phenylalanine; then R_4 is not unsubstituted phenylalanine or phenylalanine substituted with $4NO_2$ or $4N_3$; b) when R_1 is a tyrosyl residue; R_2 is D-alanine; X, Y, and Z are H; and R_4 is phenylalanine; then R_3 is not unsubstituted phenylalanine or phenylalanine substituted with $4NO_2$; c) when R_1 is a tyrosyl residue; R_2 is D-alanine; X, Y, and Z are H; and R_4 is 1'-naphthylalanine; then R_3 is not 1'-naphthylalanine or 2'-naphthylalanine; d) when R_1 is a tyrosyl residue; R_2 is D-alanine; and X, Y and Z are H; then R_3 and R_4 are not tryptophan; e) when R_1 is a tyrosyl residue; R_2 is a D-amino acid with a lower alkyl or lower thioalkyl group as a side chain; and R_4 is a neutral amino acid; then R_3 is not unsubstituted phenylalanine; and wherein said compound is not selected from the group consisting of: H-Tyr-D-Phe-Phe-Phe-NH_2; H-Tyr-D-NMePhe-Phe-Phe-NH_2; H-Tyr-D-Tic-Phe- Phe-NH_2; H-Tyr-Pro-Phe-Thr(Bzl)-NH_2 (SEQ ID NO:2); H-Tyr-Pro-Phe-Phe-NH_2 (SEQ ID NO:1); H-Tyr-Pro-Phe-Apb-NH_2; H-Tyr-Pro-Phe-App-NH_2; H-Tyr-Pro-Phe-Aph-NH_2; and H-Tyr-Pro-Apb-Phe-NH_2: wherein Apb is 2-amino-4-phenylbutanoic acid, App is 2-amino-5-phenyl pentanoic acid and Aph is 2-amino-6-phenylhexanoic acid.

On page 9 of the English translation of the specification, please amend the last paragraph which ends in page 11 to read as follows:

Specific, individual, preferred compounds of this invention are as follows:

H-Tyr-D-Ala-Phe-2'-Nal-NH_2;

H-Tyr-D-Ala-D-Phe-Phe-NH_2;

H-Tyr-D-Ala-Phe(4NO_2)-Phe(4NO_2)-NH_2;

H-Tyr-D-Ala-Phe-Tic-NH_2;

H-Tyr-D-Ala-Phe-Phe(NMe)-NH_2;

H-Tyr-D-Ala-Phe-1'-Nal-NH_2;

H-Tyr-D-Ala-Trp-Phe-NH_2;

H-Tyr-D-Ala-Phe-Trp-NH_2;

H-Tyr-∇Ala-Phe-Phe-NH_2; (SEQ ID NO:3)

∇CH_2-Tyr-D-Ala-Phe-Phe-NH_2;

H-Tyr-D-Nle-Phe-Trp-NH_2;

H-Tyr-D-Nle-Phe-Trp-NH_2;

H-Tyr-D-Nle-Phe-2'-Nal-NH_2;

H-Tyr-D-Nle-Trp-Phe-NH_2;

H-Tyr-D-Ala-Trp-2'-Nal-NH_2;

H-Tyr-D-Nle-Trp-2'-Nal-NH_2;

H-Tyr-D-Nle-Trp-Trp-NH_2;

H-Tyr-D-Nva-Phe-Phe-NH_2;

H-Tyr-D-Ser-Phe-Phe-NH_2;

H-Tyr-D-Val-Phe-Phe-NH_2;

H-Tyr-D-Leu-Phe-Phe-NH_2;

H-Tyr-D-Ile-Phe-Phe-NH_2;

H-Tyr-D-Abu-Phe-Phe-NH_2;

H-Tyr-Chl-Phe-Phe-NH_2;

H-Tyr-Cle-Phe-Phe-NH_2;

H-Tyr-D-Arg-Phe-Phe-NH_2;

H-Tyr-D-Cys-Phe-Phe-NH_2;

H-Tyr-D-Thr-Phe-Phe-NH_2;

H-DMT-D-Ser-Phe-Phe-NH_2;

H-Tyr-D-Ala-Phe-Phe-OH trifluoroacetate;

H-Tyr-D-Ala-Phe-Phg-NH_2 trifluoroacetic acid salt;

H-D-DMT-D-Ala-Phe-Phe-NH_2 trifluoroacetic acid salt;

H-Tyr-D-Ala-Phe-Hph-NH_2 trifluoroacetic acid salt;

H-Tyr-D-Ala-Phe-Cys(Bzl)-NH_2 trifluoroacetic acid salt;

H-Tyr-D-Arg-Hph-Phe-NH_2 bis-trifluoroacetic acid salt;

H-Tyr-D-Arg-Phg-Phe-NH_2 bis-trifluoro acetic acid salt;

H-Tyr-D-Ala-Phe-Phe-CH_2OH hydrochloride salt;

H-Tyr-D-Ala-Hph-Phe-NH_2 trifluoroacetic acid salt;

H-Tyr-D-Met-Phe-Phe-NH_2 trifluoroacetic acid salt;

H-Tyr-D-Arg-Phe-D-Phe-NH_2 bis-trifluoroacetic acid salt;

H-Tyr-D-Ala-Phg-Phe-NH_2 trifluoroacetic acid salt;

H-Tyr-(D)-Ala-(D)-Phg-Phe-NH_2 trifluoroacetic acid salt;

H-Tyr-D-Arg-Phe-Phe(pF)-NH_2 bis-trifluoroacetic acid salt;

H-Tyr-D-Arg-Phe-D-Phe(pF)-NH_2 ditrifluoroacetic acid salt;

H-Tyr-D-Ala-Phe-Phe(pF)-NH_2 trifluoroacetic acid salt; and

H-Tyr-D-Ala-Phe-D-Phe(pF)-NH_2 trifluoroacetic acid salt.

On page 19 of the English translation of the specification, please amend Table 1 to read as follows:

Table 1

H#	Sequence	Ki^{μ} [nM]	Ki^{δ}/Ki^{μ}	GPI(IC_{50}) [nM]	ED_{50}(PBQ) mg/k (20min)	Hot Plate mg/kg
74	H-Tyr-D-Ala-Phe-Phe-NH_2	1.53	409	3	1.4	>100
3	H-Tyr-D-Phe-Phe-Phe-NH_2	3.63	37.7	247	>20	
4	H-Tyr-Aib-Phe-Phe-NH_2			73	>20	
5	H-Tyr-D-Nle-Phe-Phe-NH_2	0.968	373	15	2.5 (5 min.)	
6	H-Tyr-Pro-Phe-Phe-NH_2 (SEQ ID NO:1)	4.10	182	15	>20	
7	H-Tyr-D-Ala-Phe-2'-Nal-NH_2	0.655	119	2	1.1 (5 min.)	
8	H-Tyr-D-Ala-2'-Nal-1'-Nal-NH_2	5.61	102	-	>20	
75	H-Tyr-D-Ala-D-Phe-Phe-NH_2	26.0	82.7	925		
76	H-Tyr-D-Ala-Phe-Phe(4-NO_2)-NH_2	0.509	129	8	4	
77	H-Tyr-D-Ala-Phe(4-NO_2)-Phe(4-NO_2)-NH_2	0.826	570	6	>20	
78	H-Tyr-D-Ala-Phe-Phe(4-N_3)-NH_2	1.49	107	50		
79	H-Tyr-D-Ala-Phe(4-NO_2)-Phe-NH_2	56.8	24.3	77		
80	H-Tyr-D-Ala-Phe-Tic-NH_2	12.7	279	-		
81	H-Tyr-D-Ala-Phe-Phe(NMe)-NH_2	22.6	215	241		
82	H-Tyr-D-Ala-Phe-1'-Nal-NH_2	0.981	174	2	>20	

n page 20 of the English translation of the specification, please amend continuation of Table 1 to read as follows:

BCH#	Sequence	Ki^{μ} [nM]	$Ki\delta/Ki^{\mu}$	GPI(IC_{50}) [nM]	ED_{50}(PBQ) m/k (20min)	Hot Plate M/k
1783	H-Tyr-D-Ala-l'-Nal-l'-Nal-NH_2	2.88	410	-	>20	
1784	H-Tyr-D-Ala-Trp-Phe-NH_2	3.57	238	20	>20	
1785	H-Tyr-D-Ala-Phe-Trp-NH_2	2.21	214	16	>20	
1786	H-Tyr-D-Ala-Trp-Trp-NH_2	0.833	783		10	
1787	H-Tyr-VAla-Phe-Phe-NH_2 (SEQ ID NO:3)				10	
2202	∇CH_2Tyr-D-Ala-Phe-Phe-NH_2				>10	
2208	H-Tyr-D-Nle-Phe-Trp-NH_2				>3	
2211	H-Tyr-D-Nle-Phe-2'-Nal-NH_2				>10	
2212	H-Tyr-D-Nle-Trp-Phe-NH_2				>10	
2213	H-Tyr-D-Ala-Trp-2'-Nal-NH_2				>5	
2214	H-Tyr-D-Nle-Trp-2'-Nal-NH_2				15	
2217	H-Tyr-D-Nle-Trp-Trp-NH_2				>5	
2462	H-Tyr-D-Nva-Phe-Phe-NH_2				2.7	>100
2463	H-Tyr-D-Ser-Phe-Phe-NH_2	2.2		13	0.5	>100
2464	H-Tyr-D-Val-Phe-Phe-NH_2				>10	
2465	H-Tyr-D-Leu-Phe-Phe-NH_2				> 10	